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Maximum Match 100%
Listing first 45 summaries
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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2: /SIDS2/gcgdata/geneseq/geneseqn/NA1981.DAT:*

3: /SIDS2/gcgdata/geneseq/geneseqn/NA1982.DAT:*

4: /SIDS2/gcgdata/geneseq/geneseqn/NA1983.DAT:*

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20: /SIDS2/gcgdata/geneseq/geneseqn/NA1995.DAT:*

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Copyright (c) 1993 - 2000 Comp
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AAC76691
AAA53484
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AAF 64883
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Probe #12901 for g
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Human CDNA clone (
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ALIGNMENTS

Human ovarian tumo	AAZ77506	20	2188		52	45	
IL-1ra	AAX02974	20	3198	1.9	52.4	44	
gen	AAD08215	22	114793	٠	53	43	
HSV L/ST region.	AAQ76213	16	12001		54.8	42	
	AAF58238	22	244		70.4	41	
Oligonucleotide D1	AAF58238	22	244		72	40	
	AAF65378	22	377		99.4	9	
secret	AAC04338	21	195		124	38	
	AA150060	22	149		149	37	
Probe #14636 for g	AAI24703	22	149		149	36	
Probe #25826 used	AAI57140	22	165		165	35	
Human gene signatu	AAT25803	16	180		٠	34	
eotide	AAF58255	22	938			33	
Oligonucleotide D2	AAF58262	22	936			32	
Oligonucleotide D2	AAF58259	22	936			31	
Oligonucleotide D1	AAF58257	22	936	٠		30	
Oligonucleotide D1	AAF58254	22	936			29	
leotide	AAF58252	22	936			28	
Oligonucleotide D1	AAF58255	22	938		•	27	
æ	AAF58262	22	936	6.4		26	
Φ	AAF58259	22	936	6.4		25	
	AAF58257	22	936	6.4		24	
	AAF58254	22	936	6.4	176.8	23	
nucleotide	AAF58252	22	936	6.4		22	
#5549 used	AAI36863	22	449			21	
#5408 for g	AAI15475	22	449			20	
#6543 used	AAI37857	22	416		٠	19	
#20179 used	AAI51493	22	201		٠	18	
#1525	AAI25319	22	201			17	
	AAI38399	22	468			16	
	AAI16129	22	468			15	
	AAF64832	22	397		11.	14	
n secret	4387	N	318	10.6	294.6	13	
Novel human polynu	\mathbf{L}	N	403	w	84.	12	

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RESULT AAA53483 ID AAA53 XX AAA53 AC AAA5 XX AAA5 XX Huma XX Huma XX Homo XX Homo XX Homo XX Homo XX Homo XX III-M XX Homo XX AAA5 PT CDS PT 04-OCT-2000 (first entry) Homo sapiens. H37; human; Cdc7 regulatory subunit; cytostatic; proliferative; cancer; anti-proliferative; replication regulator; stem cell; ss. Human H37 nucleotide sequence #1. AAA53483; AAA53483 standard; cDNA; 2780 ш BP.

Location/Qualifiers 518..2542 /*tag= a /product= "H37" /*tag=

Arai K,

Masai H;

(NISC-) JAPAN SCI & TECHNOLOGY CORP (ARAI/) ARAI K. (MASA/) MASAI H.

30-OCT-1998; 01-NOV-1999; 11-MAY-2000. WO200026250-A1

98JP-0311408 99WO-JP06076

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Query Match
Best Local Sim.
Matches 2780;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a human H37 encoding nucleotide sequence. H37 is a protein with a Cdc7 activity regulatory subunit. The invention relates to two H37 protein and nucleotide sequences. H37 proteins exhibit cytostatic, proliferative, anti-proliferative, and cell replication regulatory activities. The proteins, encoded genes and DNA fragments are useful in treating cancers and other diseases resulting from abnormal production of stem cells. Antibodies directed against one of the H37 proteins can be used to inhibit cell proliferation.
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P-PSDB; AAB03758.
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                    aaaagtattttaccttgacttaccttctgtcaccatatctgaaaaacttcaaaaggacat 720
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illarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
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1800	tcagagcccatcccccaccttcaaatgaattgagagggcttaatg	7	Дb
	ctcctgtttatttcagagcccatcccccacccttcaaatgaattgaga	74	Оу
1740	gatacaacagtgaaggagcagaatttcctgtataaagagacccaggaaactgaaaaaa	9	Дb
1740	atacaacagtgaagg	1681	Qy
1680	ggaattgcaacatattctcagaaagattgccaggaag	Q.	Db
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1620	.aaaatacagtgttggatccctttctcctgtttctgcaagtgtcctgaaaa	1561	Дb
Ġ	aaaagaataaaataca 	Ġ	γо
1560	tgactttgtggaatatgaaaaggacacctaaaa		Дb
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1500	ttgcacagagtaaccagtatcaagttgttg	4	ф
1500	caccttctaagtgagcaacacagaaactttgcacagagtaacca	4	Qy
1440	ggatattgtgaatgttgcttgcagaaatatgaagatctagaaa	ω	рь
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1380	.ccaaacagatggcgataagtatggtggaacctcaattcaactccagttga	w	дb
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1260	tataggccattttatcttcagctgaccaatatgccttttataaattattctattcaga	1201	Qγ
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1200	aaaacaagaacaggaagactcaaaaagccttttgtaaaggtggaagatatgagccaact	1141	Qy
1140	tggtagtggtgcaca	1081	Db
	ctcaagaaatcaagtacttcagtaagagatgggggcaaaagagttggtagtggtgcaca	0	Qy
0	tgaacaaaagaaaaaagagttgtattt	1021	ф
1080	attcttcatattgatgacattagatactacattgaacaaaagagaaaaaagagttgtattt	1021	Qy
1020	tatcaaatgccttgtcatggggagtaa	961	郞
1020	gaccatgattttattccttcaaatagtatattatcaaatgccttgtcatggg	6	Qy
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960	agtcaccagacacagtgtgtttaagcagaggaaaattattagttgaaaaagctatcaa	901	Qy
900	tcacctcatcccagccatgatggaagttcatt	841	DЬ
900	gaatotgcatatactgcagaaaccacttcacctcatcccagccatgatggaagttcatt	41	ОУ
840	accttgggtcgaatttctcctgtaccaagtcc	781	Db '
840	aataagaaggaagctaaatttgcacaaaccttgggtcgaatttctcctgtaccaagtcc	81	Qy
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hypotensive; dermatological; immunosuppressive; antiinflammatory; antiviral; antibacterial; antifungal; antifheumatic; antithyroid; antianaemic; gene therapy; cancer; proliferative disorder; hypertension; neurodegenerative disorder; osteoarthritis; graft vs host disease; cardiovascular disease; diabetes mellitus; hypothyroidism; SCID; AIDS; cholesterol ester storage; systemic lupus erythematosus; infection; severe combined immunodeficiency; malaria; autolimnune disorder; asthma; allergy; aplastic anaemia; nocturnal haemogloblnuria; burn; wound;
                                proliferative disorders, neurodegenerative disorders, osteoarthritis, graft vs host disease, cardiovascular disease, diabetes. mellitus, hypertension, hypothyroidism, cholesterol ester storage, systemic lupus erythematosus, severe combined immunodeficiency (SCID), AIDS, viral, bacterial or fungal infection, malaria, autoimmune disorders, asthma, allergies, aplastic anaemia, burns, wounds, bone and cartilage damage, nocturnal haemoglobinuria, antiinflammatory disease; to enhance coagulation; to inhibit thrombosis; and as a contraceptive.
                                                                                                                                                                          antiinflammatory; antibacterial; antiviral; antifungal; antirheumatic; antithyroid; and antianaemic. The sequences can be used for determining the presence of or predisposition to, or preventing or treating pathological conditions associated with an OREX-associated disorder. The nucleic acids can be used to express ORFX proteins in gene therapy vectors. The proteins and nucleic acids may be used to treat cancers,
                                                                                                                                                                                                                                                                                      antipsoriatic; antiparkinsonian; nootropic; neuroprotective; osteopathic; anticonvulsant; antiarthritic; immunosuppressant; immunostimulant; cardiant; thrombolytic; coagulant; vasotropic; antidiabetic; hypotensive; dermatological; immunosuppressive;
                                                                                                                                                                                                                                                                                                                                                            which represent the human ORFX open reading frames 1 to 3161. sequences have activities such as: cytostatic; hepatotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel nucleic acids and peptides derived from open reading useful for treating e.g. cancers, proliferative disorders, neurodegenerative disorders and cardiovascular disease -
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RESULT 4
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Probe; human; microarray;
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OS Homo sapiens.
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OS Homo sapiens.
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PN W0200157278-A2.
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PN 09-AUG-2001; 2001WO-US006-
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PR 04-PEB-2000; 2000US-01803-
PR 26-MAY-2000; 2000US-02074-
PR 30-JUN-2000; 2000US-02346-
PR 30-JUN-2000; 2000US-02363-
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Best Local Similarity
Matches 1196; Conserv
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Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ftp.wipo.int/pub/published_pct_sequences.
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2000US-0632366.
2000US-0233687.
2000US-0236359.
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Pred. No. 5.2e-258;
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09-OCT-2001 (first entry)

Probe; human; breast disease; breast cancer; development disorder; ss; inflammatory disease; proliferative breast disease; non-carcinoma tumour. probe #3618 used to measure gene expression in human breast sample

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26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0684408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234667.
27-SEP-2000; 2000US-02345359.
                                                                                                                                                                                                                                                                                                                                                                The present invention relates to novel single exon nucleic acid probes. The present sequence is one such probe. The probes are useful for measuring human gene expression in a human breast sample, where the probe hybridises at high stringency to a nucleic acid expressed in the human breast. The probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human breast, particularly those diseases with polygenic actiology. The diseases include: breast cancer disorders of development, inflammatory diseases of the breast, fibrocystic changes, proliferative breast disease and non-carcinoma tumours.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                      Novel single exon nucleic acid probe used to measuring gene expression in a human breast \boldsymbol{\cdot}
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                                       (MOLE-) MOLECULAR DYNAMICS INC.
1803 aaatgagtaataaatgttccatgttaagtacagctgaagatgacataagacagaatttta 1862
                                                                                                                                                                                                                                                                                                                          Sequence 1915 BP; 608 A; 324 C; 280 G; 703 T; 0 other;
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                          TTACAGACCCAAATGTAAATATTAAAAATAATATTTGCAATTTTCTACAGAATTGAATA 144
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RESULT AAH17747 ID AAH1 XX AAH1 XX AAH1 XX 26-0 DT 26-0 XX XX Human cDNA sequence SEQ ID NO:17368 26-JUN-2001 (first entry) AAH17747; AAH17747 standard; cDNA; 2276 6

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The present invention describes primer sets for synthesising 5602 CC full-length cDNAs defined in the specification. Where a primer set CC comprises: (a) an oligo-dT primer and an oligonucleotide comprises one of the 5602 nucleotide sequences defined in the specification, where the CC cligonucleotide comprises at least 15 nucleotides; or (b) a combination of an oligonucleotide comprising a sequence complementary to the CC complementary strand of a polynucleotide which comprises a 5'-end CC sequence and an oligonucleotide comprises a 3'-end sequence, where the CC cligonucleotide which comprises a 3'-end sequence, where the CC polynucleotide which comprises a 3'-end sequence, where the CC digonucleotide comprises a 3'-end sequence, where the CC cligonucleotide comprises a 3'-end sequence, where the CC cligonucleotide comprises at least 15 nucleotides and the combination of the 5'-end sequence/3'-end sequence is selected from those defined in CC the 5'-end sequence/3'-end sequence sets can be used in antisense therapy and CC in gene therapy. The primers are useful for synthesising polynucleotides, particularly full-length cDNAs. The primers are also useful for the CC detection and/or diagnosis of the abnormality of the proteins encoded by the full-length cDNAs. The primers allow obtaining of the full-length CC cDNAs easily without any specialised methods. AAH03166 to AAH13632 and AAH13633 to AAH13632 represent human amino acid sequences; AAB92446 to AAB95893 represent human amino acid sequences; and AAH13632 to AAH13632.
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27-AUG-1999; 99JF-0300253.
11-JAN-2000; 2000JF-0118776.
02-MAY-2000; 2000JF-0183767.
09-JUN-2000; 2000JF-0241899.
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30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
21-SEB-2000; 2000US-0234687.
27-SEB-2000; 2000US-0234635.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to human single exon nucleic acid probes (SENP). The present sequence is one such probe. The SENPs are derived from human Hela cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staying of diseases of the cervix, notably cervical cancer. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at the cervix and the content of the content of the cervix and the content of the cervix and the cervity in the cervical cancer.
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                                                                                                                                                                                                                                                                             Probe #8625 used to measure gene expression in human breast sample.
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2000US-0180312.
2000US-0207456.
2000US-0608408.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim
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                                               cacagctacctctacataaaaacaaaacaggaatgcattcttgacatttccgaacacacat
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             atgattctggtctgataacaataaacagttcacaagagcacctaactgttcaggcaaagg
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27-AUG-1999;
11-JAN-2000;
02-MAY-2000;
09-JUN-2000;
                                                                                                                                             Primer sets for synthesizing polynucleotides, particularly the full-length cDNAs defined in the specification, and for the det and/or diagnosis of the abnormality of the proteins encoded by full-length cDNAs -
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Ishii :
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99JP-0300253.
2000JP-0118776.
2000JP-0183767.
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Otsuki
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The present invention describes primer sets for synthesising 5602 full-length cDNAs defined in the specification. Where a primer set comprises: (a) an oligo-dT primer and an oligonucleotide complementary to the complementary strand of a polynucleotide which comprises one of

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cc the 5602 nucleotide sequences defined in the specification, where the colliponucleotide comprises at least 15 nucleotides; or (b) a combination cc of an oligonucleotide comprising a sequence complementary to the complementary strand of a polynucleotide which comprises a 5'-end complementary to the oligonucleotide which comprises a 5'-end sequence and an oligonucleotide comprising a sequence complementary to a cc polynucleotide which comprises a 1'-end sequence, where the combination of cc oligonucleotide comprises at least 15 nucleotides and the combination of ct the 5'-end sequence's'-end sequence's'-end sequence's and the combination of cc in gene therapy. The primer sets can be used in antisense therapy and cc in gene therapy. The primers are useful for synthesising polynucleotides, can be call-length cDNAs. The primers are also useful for the cc detection and/or diagnosis of the abnormality of the full-length cDNAs. The primers allow obtaining of the full-length cc compasses the compasses of the full-length cc compasses the compasses of the full-sequence (cc pNAs seasily without any specialised methods. ANH03165 to ANH13632 compasses to ligonucleotides, all of which are used in the exemplification cc represent oligonucleotides, all of which are used in the exemplification cc of the present invention.
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661 agaactgaatttattnccccagaagaaaacngaatttgt-tttcaccggtacagtcttta 719
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pred. No. 3.1e-152;
0; Mismatches 56;
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Sequence 685 BP; 238 A; 127 C; 125

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RESULT 1
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The present sequence is one of 3351 sequences in a library of human CC polynucleotides. The library is used to detect differentially expressed CC genes correlated with a cancerous state of a mammalian cell and can cell detect colon, prostate, breast and lung cancer. The library can be used to produce probes for detection of mRNA and to produce additional copies CC of the polynucleotides. The probes can be used for chromosome mapping of CC or antisense oligonucleotides can be generated. The polynucleotides and cc their gene products are used as genetic or biochemical markers (e.g. in CC plood or tissues) that will detect the earliest changes along the CC carcinogenests pathway and/or monitor the efficacy of therapies and CC antibodies against them can be used in pharmaceutical compositions to CC creat the cancers and proliferative disorders such as neoplasia, and dysplasia and hyperplasia.
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                                                                                                                                                                                                                                                                                                                               Library of polynucleotides for diagnosing a cancerous state of mammalian cell and detecting cancer, particularly of the colon prostate, comprises 3351 human polynucleotide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                               Williams LT, Escobedo J, Innis MA, Garcia PD, Klinger J, Kassam
Reinhard C, Randazzo F, Kennedy GC, Pot D, Lamson G, Drmanac R;
Crkenjakov R, Drmanac S, Dickson M, Liabat I, Leshkowittz D;
Kita D, Garcia V, Jones LW, Strache-Crain B;
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02-JUL-1999;
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    Williams
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                                             (CHIR ) CHIRON CORP. (HYSE-) HYSEQ INC.
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                                                                                                                                                                                                                                                                                                                                                                                             cytostatic; gene therapy; colon cancer; prostate
  LT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                        polynucleotide,
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  Escobedo
                                                                                                                                                                                                                                                                                                                                                                        lung
                                                                                                                99US-0142310
99US-0142311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA;
                                                                                                                                                                                                                                                                                                                                                                          cancer; cancer detection;
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94.9%;
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Pred. No. 1e-105;
0; Mismatches 2
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MΑ,
  Garcia
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  PD,
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Klinger
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CC The present sequence is one of 3351 sequences in a library of human CC polynucleotides. The library is used to detect differentially expressed CC genes correlated with a cancerous state of a mammalian cell and can CC detect colon, prostate, breast and lung cancer. The library can be used CC of the polynucleotides. The probes can be used for chromosome mapping of CC of the polynucleotides. The probes can be used for chromosome mapping of CC or antisense oligonucleotides can be generated. The polynucleotides and CC their gene products are used as genetic or blochemical markers (e.g. in CC blood or tissues) that will detect the earliest changes along the CC carcinogenesis pathway and/or monitor the efficacy of therapies and CC antibodies against them can be used in pharmaceutical compositions to CC treat the cancers and proliferative disorders such as neoplasia, XX dysplasia and hyperplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Crkenjakov R,
Kita D, Garci
                                                                                                                                                                                                                                                                                                                                           Claim 9; Page 915; 1046pp; English
                                                                                                                                                                                                                                                                                                                                                                                                Library of polynucleotides for diagnosing a cancerous state of mammalian cell and detecting cancer, particularly of the colon
Sequence 422 BP; 147 A; 77
                                                                                                                                                                                                                                                                                                                                                                                prostate, comprises 3351 human polynucleotide
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Drmanac S,
ia V, Jones
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Dickson M, Labat I,
LW, Strache-Crain B;
 75
 G; 123 T;
 0 other;
                                                                                                                                                                                                                                                                                                                                                                                sequences
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Best Local Similarity 100. Matches 411; Conservative Query Match Best Local 9 1023 372 312 963 252 903 192 783 12 72 aagtattttaccttgacttaccttctgtcaccatatctgaaaaacttcaaaaggacatta ttottoatattgatgacattagatactacattgaacaaaagaaaaaaagagt 1073 accatgattttattccttcaaatagtatattatcaaatgccttgtcatggggagtaaaaa 1022 agtcaccagacacagtgtgtttaagcagaggaaaattattagttgaaaaaggtatcaagg aatctgcatatactgcagaaaccacttcacctcatcccagccatgatggaagttcatta ataagaaggaagctaaatttgcacaaaccttgggtcgaatttctcctgtaccaagtccag ataagaaggaagctaaatttgcacaaaccttgggtcgaatttctcctgttaccaagtccag aggatctgggagggcgagttgaagaatttctcagcaaagatatcagttatcttatttcaa aagtattttaccttgacttaccttctgtcaccatatctgaaaaacttcaaaaaggacatta agtcaccagacacagtgtgtttaagcagaggaaaattattagttgaaaaagctatcaagg aatctgcatatactgcagaaaccacttcacctcatcccagccatgatggaagttcattta aggatctgggagggcgagttgaagaatttctcagcaaagatatcagttatcttatttcaa accatgattttattccttcaaatagtatattatcaaatgccttgtcatggggggtaaaaa ttcttcatattgatgacattagatactacattgaacaaaagaaaaaagagt 14.8%; 0; Mismatches Score 411; DB 22; pred. No. 9.2e-83; 0; Length 422; Indels 0,: Gaps 311 962 902 191 131 71 251 0

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RESULT 12
AAF64912
ID AAF64912 standard; cDNA; 403 BP.
XX
AC AAF64912;
XX
DT 09-APR-2001 (first entry)
XX
XX
DE Novel human polynucleotide, SEQ ID NO: 668.
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The present sequence is one of 3351 sequences in a library of human complying present sequence is one of 3351 sequences in a library of human complying processed sequences correlated with a cancerous state of a mammalian cell and can detect colon, prostate, breast and lung cancer. The library can be used to produce probes for detection of mRNA and to produce additional copies of the polynucleotides. The probes can be used for chromosome mapping of the polynucleotide and for detection of transcription levels. Ribozymes of antisense oligonucleotides can be generated. The polynucleotides and their gene products are used as genetic or biochemical markers (e.g. in color or tissues) that will detect the earliest changes along the carcingenesis pathway and/or monitor the efficacy of therapies and compositions to the carcingenesis pathway and/or monitor the efficacy of therapies and contibudes against them can be used in pharmaceutical compositions to treat the cancers and proliferative disorders such as neopiasia, and dysplasia and hyperplasia.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Library of polynucleotides for diagnosing a cancerous state of a mammalian cell and detecting cancer, particularly of the colon or prostate, comprises 3351 human polynucleotide sequences
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02-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 403 BP; 144 A; 82 C; 76 G; 101 T; 0 other;
                                                                                                                                                                                                                                                                                      198
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                 ctazatttgcacaaaccttgggtcgaatttctcctgtzccaagtccagaatctgcatata 854
                                                                                  ttgacttaccttctgtcaccatatctgaaaaaacttcaaaaggacattaaggatctgggag 734
                                                                                                                                                                                                                       aaactgataacaggccagaaaaatccaaatgtaagccactttggggaaaagtattttacc 674
ctamatttgcacamaccttgggtcgmatttctcctgtmccmagtccmgamtctgcmtmta
                                                                 ttgacttaccttctgtcaccatatctgaaaaacttcaaaaaggacattaaggatctgggag
                                                                                                                                                                                                     aaactgataacaggccagaaaaatccaaatgtaagccactttggggaaaagtattttacc 137
                                                                                                                                                                                                                                                                    agttccagggtggaatccaagtcaaaaatgaaaaaaacagaccatctctgaaatctctga 77
                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                       13.8%;
99.7%;
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Kennedy GC, Pot D, Lamson G, Drn
Dickson M, Labat I, Leshkowitiz
LW, Strache-Crain B;
                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                         Score 384.4; DB 22; Length 403; pred. No. 8.2e-77; 0; Mismatches 1; Indels 0;
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RESULT 1
AAA43876
ID AAA4
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Human; mouse; chicken; rat; secreted expressed sequence tag; sEST; expressed sequence tag; EST; probe; chemotactic; proliferative; immunomodulatory; haematopoletic; chemothnetic; analgesic; haemostatic; thrombolytic; antinflammatory; cytostatic; antibacterial; antifungal; antiulcer; osteopathic; neuroprotective; noctropic; antiparkinsonian; antiulcer; osteopathic; neuroprotective; noctropic; antiparkinsonian; vaccine; autoimmune disorder; multiple sclerosis; allergic condition; vaccine; autoimmune disorder; multiple sclerosis; allergic condition; insulin dependent diabetes; asthma; myelold cell deficiency; ulcer; lymphoid cell deficiency; burn; osteoparcosis; osteoarthritis; central nervous system disorder; halmelmer's disease; stroke; Parkinson's disease; Huntington's disease; coagulation disorder; haemophilla; thrombosis; inflammatory disorder; Crohn's disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tumour; infection; depression; psoriasis; ss.
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(GEMY) GENETICS INST INC.

Jacobs K, Merberg D, McCoy JM, Treacy M, LaVallie ER, Collins-Racie LA, Bowman MR;

Evans

WPI; 2000-317938/27.

Isolated polynucleotides, and encoded proteins, comprising secreted expressed sequence tags (sESTs), useful for treating various disordes such as autoimmune, infectious, and central nervous system disorders

Claim 1; Page 321; 803pp; English.

cc analysis to AAA45925 represent specifically claimed secreted expressed cc sequence tags (sESTs), isolated from human, mouse, chicken and rat claimed sources. The sESTs can have a range of activities depending on the tissues they were isolated from. The activities include: c chemotactic, proliferative; immunomodulatory; haematopoietic; c montactic; analyseic; haemostatic; thrombolytic; antiinflammatory; cc cytostatic; antibacterial; antifungal; antiviral; antiinflammatory; cc antiasthmatic; vulnerary; antilucer; osteopathic; neuroprotective; cc antiasthmatic; vulnerary; antilucer; osteopathic; neuroprotective; cc anticonvulsant; and antidepressant. The SESTs can be used for gene ct therapy and in vaccines. The SESTs are useful as probes for the sec claim of secules which correspond to the SESTs. Proteins encoded by the SESTs care useful as assays for determining biological activity and raising can be useful in assays for determining biological activity and raising can be useful assays for determining biological activity and raising can be useful assays for determining biological activity and raising can be useful assays for determining biological activity and raising can be useful assays for determining biological activity and raising can be useful assays for determining biological activity and raising can be useful assays for determining biological activity and calculations.

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WPI; 2001-091805/10

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RESULT 14
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Best Local
                   Williams LT,
Reinhard C, R
Crkenjakov R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (multiple sclerosis, insulin dependent diabetes), allergic conditions (asthma), myeloid or lymphold cell deficiencies, wounds, burns, ulcers, osteoporosis, osteoarthritis, central nervous system disorders (Alzhelmer's, Parkinson's, Huntington's disease, stroke), coagulation disorders (haemophilia, thrombosis), inflammatory disorders (Crohn's disease), tumours, bacterial, fungal or viral infections, depression and psoriasis. AAA45931 represent linker variants which are given in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1933
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                                                                                                      02-JUL-1999;
02-JUL-1999;
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           Kita
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                                                                                                                                                                                                                                                                       Novel human polynucleotide,
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                                                                   (CHIR ) CHIRON CORP
(HYSE-) HYSEQ INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ttctgatttcagtacagataatagtggatctcaaccaaaacagaagtcagatactgtgct
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                    Randazzo F,
Drmanac S,
                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 108 A; 69 C;
                                                                                                                                        2000WO-US18374
                                           Escobedo J,
                                                                                                    99US-0142310.
99US-0142311.
                                                                                                                                                                                                                                               gene therapy; colon cancer; prostate cancer;
                                                                                                                                                                                                                                                                                                                                             CDNA;
                                                                                                                                                                                                                                     cancer;
           Jones
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                   Kennedy GC,
Dickson M,
           , MT
                                                                                                                                                                                                                                     cancer detection;
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Pred. No. 9.9e
0; Mismatches
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           Strache-
                                                                                                                                                                                                                                                                       ID NO:
        Garcia PD,
Pot D, I
Labat I,
le-Crain B;
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                             PD, Klinger J,
Lamson G, Drn
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                      Leshkowitiz
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ID AAII61
XX AAII61
AC AAII61
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XX Probe
XX Probe;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1070
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                                Probe; human; microarray; gene expression; cervical epithelial cell;
                                                                                                    12-OCT-2001
                                                                                                                                      AAI16129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1130
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                 cervical cancer;
                                                                    Probe #6062
                                                                                                                                                                        AAI16129 standard;
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                                                                                                                                                                                                                                                           tctca
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                                                                  for gene
                                                                                                  (first
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                                                                                                                                                                        DNA;
                                                                                                    entry)
                                                                  expression
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Pred. No. 4.8e-38;
0; Mismatches 71;
                                                                  analysis in human
                                                                    cervical cell sample
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6.6%; Score 184.2; DB 22; Length 468;

Best Local Similarity 98.4%; Pred, No. 6.5e-32;

Matches 186; Conservative 0; Mismatches 3; Indels 0; Gaps
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26-MAY-2000; 2000US-0207456.

30-JUN-2000; 2000US-0608408.

03-AUG-2000; 2000US-0632366.

21-SEP-2000; 2000US-0234687.

27-SEP-2000; 2000US-02346359.

04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                     The present invention relates to human single exon nucleic acid probes (SENP). The present sequence is one such probe. The SENPs are derived from human HeLa cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at fig. wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                            Sequence 468 BP; 146 A; 92 C; 92 G; 138 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 25; SEQ ID No 6062; 487pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human cervical epithelial cells -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-488901/53.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MOLE-) MOLECULAR DYNAMICS INC.
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                                                        454 agtaagtct 462
                   916 agtgtgttt 924
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